### SEQUENCE LISTING

<110> BUMOL, Thomas Frank DOU, Shenshen GLASEBROOK, Andrew Lawrence GOULD, Kenneth Elliot HALE, John Edward HEUER, Josef Georg HUI, Kwan Yuk KHARITONENKOV, Alexei MIZRAHI, Jacques NA, Songqing NOBLITT, Timothy Wayne REIDY, Charles Arthur SONG, Ho Yeong WANG, Jian WU, Xiying ZUCKERMAN, Steven Harold <120> THERAPEUTIC APPLICATIONS OF MFLINT POLYPEPTIDES <130> 040902/0136 <140> US 09/280,567 <141> 1999-03-30 <150> US 60/113,407 <151> 1998-12-22 <150> US 60/112,933 <151> 1998-12-18 <150> US 60/112,703 <151> 1998-12-18 <150> US 60/112,577 <151> 1998-12-17 <150> US 60/099,643 <151> 1998-09-09 <150> US 60/086,074 <151> 1998-05-20 <150> US 60/079,856 <151> 1998-03-30 <160> 13 <170> PatentIn Ver. 2.0 <210> 1 <211> 900 <212> DNA <213> Homo sapiens <220> <221> CDS /. (900) <222> (1). <400> 1 atg agg /gcg ctg gag ggg cca ggc ctg tcg ctg ctg tgc ctg gtg ttg Met Arg/Ala Leu Glu Gly Pro Gly Leu Ser Leu Leu Cys Leu Val Leu

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GUST AND

ctg gtg cgg ctg ctg cag gcg ctg cgc gtg gcc agg atg ccc ggg ctg Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met Pro Gly Leu 275 280 285

gag cgg agc gtc cgt gag cgc ttc ctc cct gtg cac Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His 290 295 300 900

<210> 2 <211> 300 <212> PRT <213> Homo sapiens

<400> 2

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20 25 30

Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu Arg Leu Val
35 40 45

Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro Cys Arg Arg
50 55 60

Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His Tyr Thr Gln
65 70 75 80

Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val Leu Cys Gly
85 90 95

Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His Asn Arg Ala 100 105 110

Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Glu

His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly Thr Pro 130 135 140

Ser Gln Asn The Gln Cys Gln Pro Cys Pro Pro Gly Thr Phe Ser Ala 145 150 155 160

Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala 165 170 175

Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser Ser His Asp Thr Leu
180 185 190

Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala
195 200 205

Glu/Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile //210 215 220

Sér Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu Ala Pro Glu 225 230 235 240

Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu Lys 245 250 255

CAN.

Leu Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly Ala 260 Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met Prø Gly Leu 285 280 Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His <210> 3 <211> 936 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (25)..(924) <400> 3 getetecetg etecageaag gace atg agg geg etg gag ggg eea gge etg 51 Met Arg Ala Leu Glu Gly Pro Gly Leu 1 99 20 10 15 gct gta cgc gga gtg gca /aa aca ccc acc tac ccc tgg cgg gac gca Ala Val Arg Gly Val Ala/Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala : 30 gag aca ggg gag cgg étg gtg tgc gcc cag tgc ccc cca ggc acc ttt Glu Thr Gly Glu Arg/Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe 50 gtg cag cgg ccg t/gc cgc cga gac agc ccc acg acg tgt ggc ccg tgt 243 Val Gln Arg Pro/Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys 291 cca ccg cgc các tac acg cag ttc tgg aac tac ctg gag cgc tgc cgc Pro Pro Arg His Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg 75 tac tgc aa/c gtc ctc tgc ggg gag cgt gag gag gag gca cgg gct tgc 339 Tyr Cys Asn Val Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys 90 100 cac  $gc\phi'$  acc cac aac cgt gcc tgc cgc tgc cgc acc ggc ttc ttc gcg 387 His Al⁄a Thr His Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala 110 115 cac/gct ggt ttc tgc ttg gag cac gca tcg tgt cca cct ggt gcc ggc 435 His Ala Gly Phe Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly 130 gtg att gcc ccg ggc acc ccc agc cag aac acg cag tgc cag ccg tgc 483 ∜al Ile Ala Pro Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys 145

531

ccc cca qqc acc ttc tca qcc aqc tcc aqc tca gag cag tgc cag

Pro Pro Gly Thr Phe Ser Ala Ser Ser Ser Ser Glu Gln Cys Gln

			aac Asn													57/9
tct Ser	tcc Ser	tcc Ser	cat His	gac Asp 190	acc Thr	ctg Leu	tgc Cys	acc Thr	agc Ser 195	tgc Cys	act Thr	ggc Gly	ttc Phe	ccc Pro 200	ctc Leu	627
agc Ser	acc Thr	agg Arg	gta Val 205	cca Pro	gga Gly	gct Ala	gag Glu	gag Glu 210	tgt Cys	gag Glu	cgt Arg	gcc Ala	gtc Val 215	atc 11e	gac Asp	675
ttt Phe	gtg Val	gct Ala 220	ttc Phe	cag Gln	gac Asp	atc Ile	tcc Ser 225	atc Ile	aag Lys	agg Arg	ctg Leu	cag Gln 230	cag Arg	ctg Leu	ctg Leu	723
cag Gln	gcc Ala 235	ctc Leu	gag Glu	gcc Ala	ccg Pro	gag Glu 240	ggc Gly	tgg Trp	gct Ala	ccg Pro	aca Thr 248	ca Pro	agg Arg	gcg Ala	ggc Gly	771
			ttg Leu													819
			gac Asp													867
			atg Met 285													915
	gtg Val		tgat	ccto	ggc d	:c										936
<210> 4 <211> 300 <212> PRT <213> Homo sapiens																
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Ala	Leu	Pro	Ala 20	Leu	Leu	Pro	Val	Pro 25	Ala	Val	Arg	Gly	Val 30	Ala	Glu	
Thr	Pro	Thr/	Tyr	Pro	Trp	Arg	Asp 40	Ala	Glu	Thr	Gly	Glu 45	Arg	Leu	Val	
Cys	Ala 50/	Gln	Cys	Pro	Pro	Gly 55	Thr	Phe	Val	Gln	Arg 60	Pro	Cys	Arg	Arg	
Asp 65	Ser	Pro	Thr	Thr	Cys 70	Gly	Pro	Cys	Pro	Pro 75	Arg	His	Tyr	Thr	Gln 80	
Phe/	Trp	Asn	Tyr	Leu 85	Glu	Arg	Cys	Arg	Туг 90	Cys	Asn	Val	Leu	Cys 95	Gly	
G/Lu	Arg	Glu	Glu 100	Glu	Ala	Arg	Ala	Cys 105	His	Ala	Thr	His	Asn 110	Arg	Ala	

Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Gly His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly The Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr Ppé Ser Ala 150 Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala Leu Gly Leu Ala Leu Ile Val Pro Gly Ser Ser Ser His Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala , Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile 215 220 210 Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu Ala Pro Glu 235 Gly Trp Ala Pro Thr Pro Arg Ala/Gly Arg Ala Ala Leu Gln Leu Lys 250 Leu Arg Arg Arg Leu Thr Glu /Leu Leu Gly Ala Gln Asp Gly Ala Leu 270 260 265 Leu Val Arg Leu:Leu Gln Ala Leu Arg Val Ala Arg Met Pro Gly Leu 280 Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His 295 290 <210> 5 <211> 813 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1) ./. (813) <400> 5 gtg gca/gaa aca ccc acc tac ccc tgg cgg gac gca gag aca ggg gag Val Alá Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu 5 15 cgg/ctg gtg tgc gcc cag tgc ccc cca ggc acc ttt gtg cag cgg ccg Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro 20 144 rige ege ega gae age eec aeg aeg tgt gge eeg tgt eea eeg ege eae Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His tac acg cag ttc tgg aac tac ctg gag cgc tgc cgc tac tgc aac gtc Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val 55 50

240 ctc tgc ggg gag cgt gag gag gca cgg gct tgc cac gcc acc cag Leu Cys Gly Glu Arg Glu Glu Ala Arg Ala Cys His Ala Thr Hix aac cgt gcc tgc cgc tgc cgc acc ggc ttc ttc gcg cac gct ggt ttc 288 Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala 91y Phe 90 tgc ttg gag cac gca tcg tgt cca cct ggt gcc ggc gtg att gcc ccg 336 Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly Val/Ile Ala Pro 105 384 ggc acc ccc agc cag aac acg cag tgc cag ccg tgg ccc cca ggc acc Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro C/s Pro Pro Gly Thr 120 tto toa goo ago ago too ago toa gag cag tgo cag coo cac cgo aac 432 Phe Ser Ala Ser Ser Ser Ser Glu Gln Eys Gln Pro His Arg Asn 140 130 135 tgc acg gcc ctg ggc ctc gcc ctc aat /tg cca ggc tct tcc tcc cat Cys Thr Ala Leu Gly Leu Ala Leu Asn/Val Pro Gly Ser Ser Ser His 155 gac acc ctg tgc acc agc tgc act ggc ttc ccc ctc agc acc agg gta Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val 528 170 cca gga gct gag gag tgt gag cgt gcc gtc atc gac ttt gtg gct ttc 576 Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe 180; 185 cag gac atc toc atc aax agg ctg cag cgg ctg ctg cag gcc ctc gag 624 Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Gln Ala Leu Glu 200 195 gcc ccg gag ggc tg/g ggt ccg aca cca agg gcg ggc cgc gcg gcc ttg Ala Pro Glu Gly #rp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu 210 215 220 cag ctg aag ctg cgt cgg cgg ctc acg gag ctc ctg ggg gcg cag gac Gln Leu Lys Leu Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp 230 225 768 qqq qcq c£q ctq qtg cgg ctg ctg cag gcg ctg cgc gtg gcc agg atg Gly Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met 245 250 ccc ggg ctg gag cgg agc gtc cgt gag cgc ttc ctc cct gtg cac 813 Pro ¢ly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His 270 260 265 **∮**210> 6 **₹211> 271** <212> PRT <213> Homo sapiens <400> 6 Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu 10 Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro 20

Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg Hig Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asm Val Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala His/Ala Gly Phe Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Kys Pro Pro Gly Thr 115 120 Phe Ser Ala Ser Ser Ser Ser Glu Gln 🍕 Gln Pro His Arg Asn 135 Cys Thr Ala Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser Ser His 155 Asp Thr Leu Cys Thr Ser Cys Thr GL/y Phe Pro Leu Ser Thr Arg Val 170 Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile Ser Ile Lys Arg/Leu Gln Arg Leu Leu Gln Ala Leu Glu 200 Ala Pro Glu Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly Ala Leu Leu Vál Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met 250 Pro Gly Leu Gl/ Arg Ser Val Arg Glu Arg Phe Leu Pro Val His <210> 7 <211> 825 <212> DNA <213> Homo sapiens <220> CDS <221> <222₺ (1)..(813) <40/0>7gt/g gca gaa aca ccc acc tac ccc tgg cgg gac gca gag aca ggg gag

gtg gca gaa aca ccc acc tac ccc tgg cgg gac gca gag aca ggg gag 48 Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu 1 5 10 15

cgg ctg gtg tgc gcc cag tgc ccc cca ggc acc ttt gtg cag cgg ccg 9
Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro
20 25 30

tgc cgc cga gac agc ccc acg acg tgt ggc ccg tgt cca ccg cgc cac Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His 192 tac acg cag ttc tgg aac tac ctg gag cgc tgc cgc tac tgc aac/gtc Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Ash Val ctc tgc ggg gag cgt gag gag gca cgg gct tgc cac g¢c acc cac 240 Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His 70 75 65 aac egt gee tge ege tge ege ace gge tte tte geg seac get ggt tte Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe 8.5 90 tgc ttg gag cac gca tcg tgt cca cct ggt gcc/ggc gtg att gcc ccg 336 Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro 105 110 100 ggc acc ccc agc cag aac acg cag tgc cag tgc ccc cca ggc acc 384 Gly Thr Pro Ser Gln Asn Thr Gln Cys Øln Pro Cys Pro Pro Gly Thr 120 115 432 tto toa goo ago ago too ago toa gag cag tgo cag coo cac cgc aac Phe Ser Ala Ser Ser Ser Ser/Glu Gln Cys Gln Pro His Arg Asn 135 tgc acg gcc ctg ggc ctg gcc/ctc aat gtg cca ggc tct tcc tcc cat 480 Cys Thr Ala Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser His 150 155 145 gac acc ctg tgc acc agg/ tgc act ggc ttc ccc ctc agc acc agg qta 528 Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val 165 170 cca gga gct gag ga/g tgt gag cgt gcc gtc atc gac ttt gtg gct ttc 576 Pro Gly Ala Glu Gíu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe 185 624 cag gac atc tc/c atc aag agg ctg cag cgg ctg ctg cag gcc ctc gag Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Gln Ala Leu Glu 195 200 qcc ccq qaq qqc tqq qct ccq aca cca agg qcq qgc cqc qcq qcc ttq 672 Ala Pro Glu Gly Trp Ala Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu 210 cag ctg/aag ctg cgt cgg cgc ctc acg gag ctc ctg ggg gcg cag gac Gln Lex Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp 225 230 235 240 ggg/gcg ctg ctg gtg cgg ctg ctg cag gcg ctg cgc gtg gcc agg atg Gly Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met 255  $ot\!\!/cc$  ggg ctg gag cgg agc gtc cgt gag cgc ttc ctc cct gtg cac 813 ∕Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His 260 tgatcctggc cc 825

Guly P

<212> PRT <213> Homo sapiens <400> 8 Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Gl 10 Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala 🏸 s His Ala Thr His Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro 105 Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr 120 Phe Ser Ala Ser'Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn 135 Cys Thr Ala Leu Gly Leu Aļá Leu Asn Val Pro Gly Ser Ser His 150 Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val 170 Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe 185 180 Gln Asp Ile Ser/Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu 195 Ala Pro Glu Gly Trp Ala Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu 215 Gln Leu Lys Leu Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp 230 235 Gly Ala/ Leu Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met 250 Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His 265 **∮**210> 9

Suly (M)

<210> 8 <211> 271

,6211> 8 <212> PRT

<213> Artificial Sequence

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Asp Tyr Lys Asp Asp Asp Lys
<210> 10
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence / Primer
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tagggctgat caaggatggg cttctggact tggg/cgccc ctccgcaggc ggaccgggg 59
<210> 11
<211> 66
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
agggggggg ccgctgatca *cacttgtcg tcgtcgtcct tgtagtcgtg cacagggagg 60
                                                                   66
aagcgc
<210> 12
<211> 37
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 12
                                                                   37
gaagatct/c tttgatcaag gatgggcttc tggactt
<210>/13
<211 37
<212 > DNA
<21/3> Artificial Sequence
<£20>
(223> Description of Artificial Sequence: Primer
                                                                   37
ggactagtcc tgatcatcac ttgtcgtcgt cgtcctt
```

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# FIG. 2

									J							
geoctecetg etceageaag gace atg agg geg etg gag ggg eea gge etg Met Arg Ala Leu Glu Gly Pro Gly Leu 1 5														51		
tcg Ser 10	ctg Leu	ctg Deu	tgc Cys	ctg Leu	gtg Val 15	ttg Leu	gcg Ala	ctg Leu	cct Pro	gcc Ala 20	ctg Leu	ctg Leu	ccg Pro	gtg Val	ccg Pro 25	99
		cgc Arg														147
		ggg Gly														195
		cgg Arg 60														243
cca Pro	ccg Pro 75	cgc Arg	cac His	tac Tyr	acg Thr	cag Gln 80	ttc Phe	tgg Trp	aac Asn	tac Tyr	ctg Leu 85	gag Glu	cgc Arg	tgc Cys	cgc Arg	291
tac Tyr 90	tgc Cys	aac Asn	gtc Val	ctc Leu	tgc Cys 95	Gly 999	gag Glu	egt	gag Glu	gag Glu 100	gag Glu	gca Ala	cgg Arg	gct Ala	tgc Cys 105	339
		acc Thr														387
cac His	gct Ala	ggt Gly	ttc Phe 125	tgc Cys	ttg Leu	gag Glu	cac His	gca Ala 130	tcg Ser	tgt Cys	cca Pro	cct Pro	ggt Gly 135	gcc Ala	ggc Gly	435
		gcc Ala 140														483
		ggc Gly														531
		cgc Arg		Cys		Ala			Leu		Leu					579
tct Ser	tcc Ser	tcc Ser	cat His	gac Asp 190	acc Thr	ctg Leu	tgc Cys	acc Thr	agc Ser 195	tgc Cys	act Thr	ggc Gly	ttc Phe	ccc Pro 200	ctc Leu	627
		agg Arg														675
		gct Ala 220														723

FIG. 2 (cont'd.)															
cag gcc Gln Ala 235															771
cgc gcg Arg Ala 250 '															819
ggg gcg Gly Ala		Asp (													867
gtg gcc Val Ala	Arg M														915
cct gtg Pro Val		tgat	cctg	igc c	:c \	1	1			ı					936

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## FIG. 4

					ccc Pro 5												48
	cgg Arg	ctg Leu	gtg Val	tgc Cys 20	gcc Ala	cag Gln	tgc Cys	ccc Pro	cca Pro 25	ggc Gly	acc Thr	ttt Phe	gtg Val	cag Gln 30	cgg Arg	ccg Pro	96
	tgc Cys	cgc Arg	cga Arg 35	gac Asp	agc Ser	ccc Pro	acg Thr	acg Thr 40	tgt Cys	ggc Gly	ccg Pro	tgt Cys	cca Pro 45	ccg Pro	cgc Arg	cac His	144
	tac Tyr	acg Thr 50	cag Gln	ttc Phe	tgg Trp	aac Asn	tac Tyr 55	ctg Leu	gag Glu	cgc Arg	tgc Cys	cgc Arg 60	tac Tyr	tgc Cys	aac Asn	gtc Val	192
					cgt Arg												240
					cgc Arg 85												288
	tgc Cys	ttg Leu	gag Glu	cac His 100	gca Ala	tcg Ser	tgt Cys	cca Pro	cot Pro 105	ggt Gly	gcc Ala	ggc Gly	gtg Val	att Ile 110	gcc Ala	ccg Pro	336
					cag Gln												384
					agc Ser												432
	tgc Cys 145	acg Thr	gcc Ala	ctg Leu	ggc Gly	ctg Leu 150	gcc Ala	ctc Leu	aat Asn	gtg Val	cca Pro 155	Gl <sup>X</sup>	tct Ser	tcc Ser	tcc Ser	cat His 160	480
					acc Thr 165												528
			Ala	Glu	gag Glu	Cys	Glu	Arq		Val			Phe				576
	cag Gln	gac Asp	atc Ile 195	tcc Ser	atc Ile	aag Lys	agg Arg	ctg Leu 200	cag Gln	cgg Arg	ctg Leu	ctg Leu	cag Gln 205	gcc Ala	ctc Deu	gag Glu	624
	gcc Ala	ccg Pro 210	gag Glu	ggc Gly	tgg Trp	gct Ala	ccg Pro 215	aca Thr	cca Pro	agg Arg	gcg Ala	ggc Gly 220	cgc Arg	gcg Ala	gcc Ala	ttg Deu	672
					cgt Arg												720

# X-12915 G2 CONT.

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FIG. 4 (cont'd.)																
gly aaa	gcg Ala	ctg Leu	ctg Leu	gtg Val 245	cgg Arg	ctg Leu	ctg Leu	cag Gln	gcg Ala 250	ctg Leu	cgc Arg	gtg Val	gcc Ala	agg Arg 255	atg Met	768
ccc Pro	ggg ggg	ctg Leu	gag Glu 260	cgg Arg	agc Ser	gtc	cgt Arg	gag Glu 265	cgc Arg	ttc Phe	ctc Leu	cct Pro	gtg Val 270	cac His		813
tgatcctggc cc																825